HIV2/SIV Sequence Summaries

LOCUS	COMMON	ACCESSION	LENGTH	REGION	FIRST AUTHOR	REFERENCE
HIV2MDS	MDS V-2 from Guinea-Bissau	Z48731	9525	complete genome	Becker,M.	Unpublished
HIV2U22047		U22047	10172	complete genome		Unpublished
SIU10899	bab2010	U10899	673	gag	Jin,M.J.	JVI 68: 8454 (1994) ne env sequence from this
HIV2U38294 Comment: Par	UC3 rt of a set of HIV2 env se	U38294 equences, access	534 ion numbers	env s U38294–U38303.	Barnett,S.	Unpublished
HIV2ENV1 Comment: Ap	CI171 parently an HIV-2 from	X78505 Ivory Coast. Sec	453 e also X7850	env 06–X78507 and X7	Janssens,W. /8510–X78511.	Unpublished
SIU19595 Comment: Th	181-01 is study reports serial p	U19595 assaging of SIV	1038 in microgli	env a, with resulting d	Lane, T.E. ivergence from the	Virology 212: 458 (1995) original stock. Macaque
HIV2GB122A Comment: Par	GB122A rt of a study of HIV infec	L10637 ction of macaque	1044 e nemestrina	env a. See also L10638.	Otten,R.A.	AIDS 8: 297 (1994)
SIVENV4 Comment: So	SMCI2 oty mangabey sample fro	X78508 om Ivory Coast;	453 see also X7	env 8509.	Peeters,M.	ARHR 10: 1289 (1994)
SIVENV025 Comment: no	20-188-2-08 information available at	Z31566 this time. Presu	322 mably maca	env ique viral sequence	Pelletier,E.	Unpublished
	MneCL8 PBwk4 1–10 neCL8 is a macrophage- uences, however, are gen	tropic virus that				Virology 207: 528 (1995) animals progress toward -U09098.
		U24287 were studied of	102 over time in	env five HIV-infected	Sankale,J.L. individuals from S	ARHR 11: 617 (1995) Genegal and Cape Verde.
SIU18039 Comment: Ac 69:1367, 1995		U18019 9–U18051. Pres	1578 umably mac	env 239 derived from th	Zhu,G.W. ne brain of an SIV-e	J. Neurovirology (in press, ncephalitic macaque (JVI
SIVENVW49 macC8 X86732 1038 env, nef Whatmore, A.M. JVI 69: 5117 Comment: Nef deletion in macaque virus C8 can be repaired in vivo, with a reversion to virulence. Accession numbers X86724–X86732 and X90850–X90854.						